
 W O R L D (TM)

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Msearch n.a. - n.a. database search, using Smith-Waterman algorithm

Run on: Wed Sep 2 19:23:41 1998; MasPar time 50.71 Seconds
 673.453 Million cell updates/sec

ular output not generated.

Title: >US-09-052-855A-8
 Description: (1:251) from US09052855A.seq
 Perfect Score: 251
 N.A. Sequence: 1 GGGGGGTGACGTGCTGTG.....GATGAGCGCTGATCGCC 251
 Comp: CCCCCACGTGACACGACGACC.....CTACTCCGACGTACGGGG

Scoring table: TABLE default

Gap 6

Mmatch STD : Dbase 0; Query 0

Searched: 188442 segs, 68026449 bases x 2

Post-processing: Minimum Match 08

Listing first 45 summaries

Database:

n-geneseq32
 1:part1 2:part2 3:part3 4:part4 5:part5 6:part6 7:part7
 8:part8 9:part9 10:part10 11:part11 12:part12 13:part13
 14:part14 15:part15 16:part16 17:part17 18:part18
 19:part19 20:part20 21:part21 22:part22 23:part23
 24:part24 25:part25 26:part26 27:part27 28:part28
 29:part29 30:part30 31:part31 32:part32 33:part33
 34:part34 35:part35 36:part36 37:part37 38:part38
 39:part39 40:part40

Statistics: Mean 7.773; Variance 4.608; scale 1.687

Pred. No. is the number of results predicted by chance to have a
 score greater than or equal to the score of the result being printed,
 and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description	Pred. No.
1	251	100.0	548	40	Polynucleotide sequen	5.18e-156
2	249	99.2	548	27	Human colon specific	1.37e-154
3	41	16.3	91	9	Oligonucleotide probe	1.27e-11
4	36	14.3	91	9	Oligonucleotide probe	1.24e-08
5	35	13.9	204	1	Base substituted E.co	4.80e-08
6	32	12.7	114	12	Generic DNA sequence	2.62e-06
7	32	12.7	114	12	Generic DNA sequence	2.62e-06
8	32	12.7	114	12	Generic DNA sequence	2.62e-06
9	32	12.7	178	32	Human endothelin-1 an	2.62e-06
10	32	12.7	204	1	Base substituted E.co	2.62e-06
11	30	12.0	114	12	Generic DNA sequence	3.56e-05
12	30	12.0	114	12	Generic DNA sequence	3.56e-05
13	26	10.4	114	12	Generic DNA sequence	5.61e-03

C	14	26	10.4	114	12	070470	Generic DNA sequence	5.61e-03
C	15	25	10.0	82	21	T13610	DC43 TSAR library gen	1.91e-02
C	16	24	9.6	114	12	070467	Generic DNA sequence	6.40e-02
C	17	24	9.6	114	12	070465	Generic DNA sequence	6.40e-02
C	18	24	9.6	178	32	T76405	Human endothelin-1 an	6.40e-02
C	19	23	9.2	69	21	T13583	TSAR-9 library genera	2.10e-01
C	20	23	9.2	75	21	T13612	DC43 TSAR library gen	2.10e-01
C	21	23	9.2	114	12	070468	Generic DNA sequence	2.10e-01
C	22	23	9.2	114	12	070466	Generic DNA sequence	2.10e-01
C	23	23	9.2	130	32	T76152	Human vascular cell a	2.10e-01
C	24	23	9.2	168	32	T76270	Human MDCK antisense	2.10e-01
C	25	23	9.2	172	32	T76363	Human Interleukin 8 a	2.10e-01
C	26	23	9.2	1473	30	T64548	NMDP-specific guttama	2.10e-01
C	27	23	9.2	1506	30	T64547	NMDP-specific guttama	2.10e-01
C	28	23	9.2	1969	30	T64531	Glutamate dehydrogena	2.10e-01
C	29	23	9.2	2096	30	T64543	NMDP-specific guttama	2.10e-01
C	30	23	9.2	2099	30	T64530	NMDP-specific guttama	2.10e-01
C	31	23	9.2	2137	30	T64542	NMDP-specific guttama	2.10e-01
C	32	23	9.2	2140	30	T64529	NMDP-specific guttama	2.10e-01
C	33	22	8.8	36	2	Q11195	Ballast Constituent C	6.75e-01
C	34	22	8.8	39	7	Q51787	Mixed oligonucleotide	6.75e-01
C	35	22	8.8	66	21	T13585	TSAR-9 library genera	6.75e-01
C	36	22	8.8	74	21	T13613	DC43 TSAR library gen	6.75e-01
C	37	22	8.8	81	21	T13611	DC43 TSAR library gen	6.75e-01
C	38	22	8.8	114	12	070471	Generic DNA sequence	6.75e-01
C	39	22	8.8	114	12	070473	Hydroxylamine oxidore	6.75e-01
C	40	22	8.8	5105	13	Q78177	TSAR-9 library genera	2.12e-00
C	41	21	8.4	65	21	T13586	TSAR-9 library genera	2.12e-00
C	42	21	8.4	68	21	T13584	Human ITS antisense	2.12e-00
C	43	21	8.4	89	32	T76219	MS-associated retrovi	2.12e-00
C	44	21	8.4	111	16	T02821	Thn-GPA1 fusion gene.	2.12e-00
C	45	21	8.4	2910	31	T62461		

ALIGNMENTS

RESULT 1
 ID V16671 standard; cDNA: 548 BP.
 AC V16671;
 DT 22-JUN-1998 (first entry)
 DE Polynucleotide sequence of a colon-specific gene.
 KW Colon-specific gene; probe; detection; expression; human;
 OS Homo sapiens.
 FH Key Location/Qualifiers
 FT CDS 1..406
 FN US5733748-A.
 PD 31-MAR-1998.
 PF 06-JUN-1995; 469667.
 PR 06-JUN-1995; US-469667.
 PA (HUMA-) HUMAN GENOME SCI INC.
 PI Rosen C, Yu G;
 DR WPI: 98-229823/20.
 DR P-PSDB: W46878.
 PT Colon-specific nucleic acids - useful as probes for detecting colon
 cancer micrometastases
 PS Claim 15: Fig 4: 51p; English.
 CC V16668-81 represent polynucleotide sequences of partial or full-length
 CC cDNA clones of colon-specific genes. The polynucleotides can be used
 CC as probes to detect expression of the corresponding human genes, e.g. in
 CC diagnostic assays for detecting micrometastases of colon cancer.
 CC Recombinant cells containing the polynucleotides can be used to
 CC produce the polypeptides. In order that antibodies can be raised and
 CC used in further screening or diagnostics. 173 G: 107 T:
 SQ Sequence 548 BP: 137 A: 128 C: 173 G: 107 T:
 Query Match 100.0%; Score 251; DB 40; Length 548;
 Best Local Similarity 100.0%; Pred. No. 5.18e-156;
 Matches 251; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
 Db 144 ggggggtgcactgtctcgtgtgctcttcagcagatccaggacaaatgaagatga 203
 |||||||

Dy	1	G6GGGCTGCACACTGTCTGTCTGTGGCCTTCATCGACAGATCCAGGAGCAAAATGAACATATA	60
Db	204	aagcagaagaactcttctctgacctggagggttcctacgcacaagaactcggtctccggga	263
Oy	61	AAGGAGAAACTTCTCTGACTTTGGGAGATTCTTAGCAAACAATCGGTCCGGGA	120
Db	264	cagctgggtctcatagagaccagaagactcaggggttaaaagccctttgagcagtctt	323
Oy	121	CAGCTGGGTCTTATAGAGACCAGAACCTCAGGGGTAAAAGCCCTTTGAGAGATTCTT	180
Db	324	aaagaaaagcccagagaccacaacaatacgaaggatggccagagctctgtgagattgaggg	383
Oy	181	AAAGACAAGCCCAGACACAAACAAATACGAGGATGGCCAGAGCTCTGAGATGAGGG	240
Db	384	ctgcagtcccc	394
Oy	241	CTGCATGCCCC	251
ID	745883	standard; cDNA; 548 BP.	
AC	T45883;		
DT	13-MAR-1997	(first entry)	
DE	Human colon specific gene CS64 cDNA partial clone.		
KW	Colon specific gene; CSG4; colon cancer; metastasis; diagnosis;		
OS	Homo sapiens.		
MH	Key	Location/Qualifiers	
FT	cds	1..408	
FT		/tag= a	
PD	WO9639419-A1.		
PD	12-DEC-1996.		
PF	06-JUN-1995; WO7289.		
PR	06-JUN-1995; WO-U07289.		
PA	(HUMA-) HUMAN GENOME SCI INC.		
PI	Rosen CA, Xu G:		
DR	WPI: 97-043054/04.		
P-PSDI	P-PSDI: W06547.		
PT	Human colon specific genes and their expression products - detection		
PT	of which, in non-colon tissue samples, can be used as indication of		
PT	colon cancer metastasis		
PS	Claim 1; Fig 4; 60pp; English.		
CC	13 cDNA clones (T45880-92), most of them partial clones, correspond		
CC	to human colon specific genes, designated CSG1, CSG2, etc., that		
CC	are primarily expressed in tissues derived from the colon. CSG7		
CC	and CSG10 show reduced expression in colon cancer cells as compared		
CC	to that in normal cells; the remaining genes are overexpressed in		
CC	colon cancer. The partial cDNA sequences can be used to isolate		
CC	full-length clones and genomic clones including the complete gene.		
CC	CSG nucleic acids can be used to produce CSG polypeptides (see also		
CC	W06545-53) in transformed host cells, as probes to detect disorders		
CC	of the colon, partic. colon cancer and colon cancer metastasis, and		
CC	in gene therapy.		
SCQ	Sequence	548 BP; 137 A; 129 C; 172 G; 107 T;	
Query Match	99.2%; Score 249; DB 27; Length 548;		
Best Local Similarity	99.6%; Pred. No. 1.37e-154;		
Matches	250; Conservative 0; Mismatches 1; Indels 0; Gaps 0;		
Db	144	gggggggtgcaactggtgtgtgtgtgctctctaagcagatccagggagccaataatgaaacgatga	203
Oy	1	GGGGGGTGTCACGTGGTGTGGGCTCTCTACGACATCCAGGAGACAAAAGAACGATGA	60
Db	204	aagcagaagaactcttctctgacctggagggttcctacgcacaagaactcggtctccggga	263
Oy	61	AAGCAGAAACTTCTCTGACTTTGGGAGATTCTTAGCAAACAATCGGTCCGGGA	120
Db	264	cagctgggtctcatagagaccagaagactcaggggttaaaagccctttgagcagtctt	323
Oy	121	CAGCTGGGTCTTATAGAGACCAGAACCTCAGGGGTAAAAGCCCTTTGAGAGATTCTT	180
Db	324	aaagaaaagcccagagaccacaacaatacgaaggatggccagagctctgtgagattgaggg	383
Oy	181	AAAGACAAGCCCAGACACAAACAAATACGAGGATGGCCAGAGCTCTGAGATGAGGG	240
Db	384	ctgcagtcccc	394
Oy	241	CTGCATGCCCC	251
ID	745883	standard; cDNA; 548 BP.	
AC	T45883;		
DT	13-MAR-1997	(first entry)	
DE	Human colon specific gene CS64 cDNA partial clone.		
KW	Colon specific gene; CSG4; colon cancer; metastasis; diagnosis;		
OS	Homo sapiens.		
MH	Key	Location/Qualifiers	
FT	cds	1..408	
FT		/tag= a	
PD	WO9639419-A1.		
PD	12-DEC-1996.		
PF	06-JUN-1995; WO7289.		
PR	06-JUN-1995; WO-U07289.		
PA	(HUMA-) HUMAN GENOME SCI INC.		
PI	Rosen CA, Xu G:		
DR	WPI: 97-043054/04.		
P-PSDI	P-PSDI: W06547.		
PT	Human colon specific genes and their expression products - detection		
PT	of which, in non-colon tissue samples, can be used as indication of		
PT	colon cancer metastasis		
PS	Claim 1; Fig 4; 60pp; English.		
CC	13 cDNA clones (T45880-92), most of them partial clones, correspond		
CC	to human colon specific genes, designated CSG1, CSG2, etc., that		
CC	are primarily expressed in tissues derived from the colon. CSG7		
CC	and CSG10 show reduced expression in colon cancer cells as compared		
CC	to that in normal cells; the remaining genes are overexpressed in		
CC	colon cancer. The partial cDNA sequences can be used to isolate		
CC	full-length clones and genomic clones including the complete gene.		
CC	CSG nucleic acids can be used to produce CSG polypeptides (see also		
CC	W06545-53) in transformed host cells, as probes to detect disorders		
CC	of the colon, partic. colon cancer and colon cancer metastasis, and		
CC	in gene therapy.		
SCQ	Sequence	548 BP; 137 A; 129 C; 172 G; 107 T;	
Query Match	99.2%; Score 249; DB 27; Length 548;		
Best Local Similarity	99.6%; Pred. No. 1.37e-154;		
Matches	250; Conservative 0; Mismatches 1; Indels 0; Gaps 0;		
Dy	324	aaagaaaagcccagagaccacaacaatacgaaggatggccagagctctgtgagattgaggg	383
Oy	181	AAAGACAAGCCCAGACACAAACAAATACGAGGATGGCCAGAGCTCTGAGATGAGGG	240
Db	384	ctgcagtcccc	394
Oy	241	CTGCATGCCCC	251
ID	745883	standard; cDNA; 548 BP.	
AC	T45883;		
DT	13-MAR-1997	(first entry)	
DE	Human colon specific gene CS64 cDNA partial clone.		
KW	Colon specific gene; CSG4; colon cancer; metastasis; diagnosis;		
OS	Homo		

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Oy 181 AAGGACAGGCCAGACACAAACAAATACGAGGGATGCCAGAGCTCTGGAGATGGAGGG 240
Db 384 ctgcagccccc 394
Oy 241 CTCGATGCCCC 251

RESULT 3
ID Q51746 standard; cDNA; 91 BP.
AC Q51746;
DT 31-MAY-1994 (first entry)
DE Oligonucleotide probe MK14-A
KW Oligonucleotide; DNA probe; mycobacteria; disease diagnosis;
OS Synthetic.
PN EP-571911-A.
PF 01-DEC-1993.
PR 24-MAY-1993: 108325.
PR 26-MAY-1992: US-889651.
PA (BECTON ) BECTON DICKINSON CO.
PI Shank DD, Spears PA;
PI WPI: 93-378844/48.
PT New oligo:nucleotide probes specific for Mycobacteria - used for
PT detection and amplification of Mycobacteria nucleic acid in
PT samples
PS Claim 3; Page 14; 23pp; English.
CC Oligonucleotide probe MK14-A consists of nucleotides 5-95 of MK14
CC (Q51735). It hybridized to all spp. of mycobacteria tested, but
CC cross reacted to a few non-mycobacterial spp. The probe may
CC be useful as an initial screen for mycobacterial infection.
CC See also Q51735-45 and Q51747-59.
SQ Sequence 91 BP; 5 A; 17 C; 15 G; 4 T;

Query Match 16.3%; Score 41; DB 9; Length 91;
Best Local Similarity 0.0%; Pred. No. 1,27e-11;
Matches 0; Conservative 45; Mismatches 4; Indels 0; Gaps 0;

Db 12 svhsyyvvvshhsbhsvhvnhvsvvvvvvvvvvvvvvvvvvvvvvvvvvvvvvv 60
Oy 16 GCTGTGGCTCTCCTACGACGATCCAGGAGCAAAATGACATGAAAGC 64

RESULT 4
ID Q51746 standard; cDNA; 91 BP.
AC Q51746;
DT 31-MAY-1994 (first entry)
DE Oligonucleotide probe MK14-A
KW Oligonucleotide; DNA probe; mycobacteria; disease diagnosis;
OS Synthetic.
PN EP-571911-A.
PF 01-DEC-1993.
PF 24-MAY-1993: 108325.
PR 26-MAY-1992: US-889651.
PA (BECTON ) BECTON DICKINSON CO.
PI Shank DD, Spears PA;
PI WPI: 93-378844/48.
PT New oligo:nucleotide probes specific for Mycobacteria - used for
PT detection and amplification of Mycobacteria nucleic acid in
PT samples
PS Claim 3; Page 14; 23pp; English.
CC Oligonucleotide probe MK14-A consists of nucleotides 5-95 of MK14
CC (Q51735). It hybridized to all spp. of mycobacteria tested, but
CC cross reacted to a few non-mycobacterial spp. The probe may
CC be useful as an initial screen for mycobacterial infection.
CC See also Q51735-45 and Q51747-59.
SQ Sequence 91 BP; 5 A; 17 C; 15 G; 4 T;

Query Match 14.3%; Score 36; DB 9; Length 91;
Best Local Similarity 0.0%; Pred. No. 1.24e-08;
Matches 0; Conservative 43; Mismatches 7; Indels 0; Gaps 0;

Db 11 ssvhsyyvvvshhsbhsvhvnhvsvvvvvvvvvvvvvvvvvvvvvvvvvvvvvvv 60

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MPearch_n n.a. - n.a. database search, using Smith-Waterman algorithm

Run on: Wed Sep 2 19:38:28 1998; MapPar time 41.52 Seconds 612.759 Million cell updates/sec

aler output not generated.

Title: >US-09-052-855A-9
Description: (1-187) from US09052855A.seq
Perfect Score: 184
N.A. Sequence: 1 GGCATGCGCCAGACTCTGG.....CTGCAATGCTTGAAGNNG 187
Comp: CCTACCGGCTCTGAGACAC.....GAGCTTCAGAACTTCNNC

Scoring table: TABLE default

Match STD : Dbase 0; Query 0

Searched: 188442 segs, 68026449 bases x 2

Post-Processing: Minimum Match 0%

Listing first 45 summaries

Database:

n-geneseq32
1:part1 2:part2 3:part3 4:part4 5:part5 6:part6 7:part7
8:part8 9:part9 10:part10 11:part11 12:part12 13:part13
14:part14 15:part15 16:part16 17:part17 18:part18
19:part19 20:part20 21:part21 22:part22 23:part23
24:part24 25:part25 26:part26 27:part27 28:part28
29:part29 30:part30 31:part31 32:part32 33:part33
34:part34 35:part35 36:part36 37:part37 38:part38
39:part39 40:part40

Statistics: Mean 7.525; Variance 5.343; scale 1.408

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description	Pred. No.
1	152	82.6	548	27	Human colon specific	7.00e-72
2	152	82.6	548	40	Polynucleotide sequen	7.00e-72
3	42	22.8	91	9	Oligonucleotide probe	6.57e-10
4	38	20.7	204	1	Base substituted E.co	7.17e-08
5	37	20.1	204	1	Base substituted E.co	2.28e-07
6	35	19.0	91	9	Oligonucleotide probe	2.27e-06
7	31	16.8	114	12	Generic DNA sequence	2.04e-04
8	30	16.3	114	12	Generic DNA sequence	6.13e-04
9	28	15.2	114	12	Generic DNA sequence	5.39e-03
10	28	15.2	114	12	Generic DNA sequence	5.39e-03
11	28	15.2	114	12	Generic DNA sequence	5.39e-03
12	28	15.2	114	12	Generic DNA sequence	5.39e-03
13	28	15.2	114	12	Generic DNA sequence	5.39e-03

C	14	28	15.2	114	12	070465	Generic DNA sequence	5.39e-03
C	15	28	15.2	114	12	070467	Generic DNA sequence	5.39e-03
C	16	27	14.7	114	12	070467	Generic DNA sequence	1.57e-02
C	17	27	14.7	130	32	T76452	Chymase antisense o11	1.57e-02
C	18	26	14.1	114	12	070468	Generic DNA sequence	4.53e-02
C	19	26	14.1	178	32	T76405	Human endothelin-1 an	4.53e-02
C	20	26	14.1	178	32	T76405	Human endothelin-1 an	4.53e-02
C	21	26	14.1	1446	38	T97343	Calcitonin gene relat	4.53e-02
C	22	25	13.6	190	32	T76452	Chymase antisense o11	1.29e-01
C	23	24	13.0	69	21	T13583	TSAR-9 library genera	3.61e-01
C	24	23	12.5	114	12	070470	Generic DNA sequence	9.87e-01
C	25	23	12.5	114	12	070472	Generic DNA sequence	9.87e-01
C	26	22	12.5	114	12	070473	Generic DNA sequence	9.97e-01
C	27	22	12.0	66	21	T13585	TSAR-9 library genera	2.71e+00
C	28	22	12.0	75	21	T13612	DC43 TSAR library gen	2.71e+00
C	29	22	12.0	82	21	T13610	DC43 TSAR library gen	2.71e+00
C	30	22	12.0	114	12	070471	Generic DNA sequence	2.71e+00
C	31	22	12.0	2478	31	T67287	Soluble starch synthn	2.71e+00
C	32	22	12.0	3871	2	N71302	HSV-1 gB and surround	2.71e+00
C	33	22	12.0	6803	33	T85473	Genomic hner sequence	2.71e+00
C	34	22	12.0	6803	40	V16305	Genomic DNA encoding	2.71e+00
C	35	21	11.4	54	25	T42303	Maize ribosomal inh1b	7.22e+00
C	36	21	11.4	65	21	T13586	TSAR-9 library genera	7.22e+00
C	37	21	11.4	68	21	T13584	TSAR-9 library genera	7.22e+00
C	38	21	11.4	81	21	T13611	DC43 TSAR library gen	7.22e+00
C	39	21	11.4	89	32	T76219	Human IL5 antisense o	7.22e+00
C	40	21	11.4	114	12	070472	Generic DNA sequence	7.22e+00
C	41	21	11.4	457	31	T45133	Wilms' tumour gene ex	7.22e+00
C	42	21	11.4	857	13	079462	Human S86/S109 Flt3 l	7.22e+00
C	43	21	11.4	968	13	079079	Human Flt-3 ligand cd	7.22e+00
C	44	21	11.4	1680	16	T02461	Wilms' tumour WT1 CDN	7.22e+00
C	45	21	11.4	1934	25	T42302	Maize ribosomal inh1b	7.22e+00

ALIGNMENTS

RESULT	1	ALIGNMENTS
ID	T45883	standard; CDNA; 548 BP.
AC	T45883:	
DT	13-MAR-1997	(first entry)
DE	Human colon specific gene CSG4 CDNA partial clone.	
KW	Colon specific gene; CSG4; colon cancer; metastasis; diagnosis;	
KM	gene therapy; ss.	
OS	Homo sapiens.	
FH	Key	Location/Qualifiers
FT	Cds	1..408
FT		/*tag- a
PN	W0639419-A1.	
PD	12-DEC-1996.	
PF	06-JUN-1995; U07289.	
PR	06-JUN-1995; WO-U07289.	
PA	(HUMA-) HUMAN GENOME SCI INC.	
PI	Rosen CA, Yu G;	
DR	WPI: 97-043054/04.	
DR	P-PSDB: W06547.	
PT	Human colon specific genes and their expression products - detection of which, in non-colon tissue samples, can be used as indication of	
PT	Colon cancer metastasis	
PS	Claim 1: Fig 4; 60pp; English.	
CC	13 CDNA clones (T45880-92), most of them partial clones, correspond	
CC	to human colon specific genes, designated CSG1, CSG2, etc., that	
CC	are primarily expressed in tissues derived from the colon. CSG	
CC	and CSG10 show reduced expression in colon cancer cells as compared	
CC	to that in normal cells; the remaining genes are overexpressed in	
CC	colon cancer. The partial CDNA sequences can be used to isolate	
CC	full-length clones and genomic clones including the complete gene.	
CC	CSG nucleic acids can be used to produce CSG polypeptides (see also	
CC	W06545-53) in transformed host cells, as probes to detect disorders	
CC	of the colon, partic. colon cancer and colon cancer metastasis, and	
CC	in gene therapy.	
SQ	Sequence 548 BP: 137 A: 129 C: 172 G: 107 T:	
Query Match	82.6%; Score 152; DB 27; Length 548;	

